

002220" 6FE0E560

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I C-----C-C-C-----CC---C-G-C
*

II G-G-C---DC-G-CCA-----C-P---G-C-----RC-C-G-L-C
* *

Fig. 1

1	6C6GTG6C66C	✓	phdtk-3	2.1	✓
1	✓	pcdtk-3	2.2	✓
1	✓	pmdtk-2	2.3	✓
1	✓	phdtk-2	2.4	✓
1	6ACCCACGCTCCGTGCTTGTGCTTCC	✓	pmdtk-1	2.5	✓
1	TGGCCCCCGCACGCAAAATTC	✓	phdtk-1	2.6	✓
1	6ACAGTCCGAGCCGCGCTGACAGCATCAAGGACTTATCTTGGAGGACTTGTGAAATCT	✓	pRNdtk-1	2.7	✓
13	6CTCTAGAAATAGTGGATCCCCCGGCTGACAGGAATTCGGCACGAGCGGCTGCGGCGCGCA6	✓	phdtk-3		
1	✓	pcdtk-3		
1	✓	pmdtk-2		
1	✓	phdtk-2		
34	GA6ATGATGGTTGTGTGTGACACGCGCA6CTGTCCGGTTCCTGGCCGTGTTACAAATGATG	✓	pmdtk-1		
23	GGCAGCgAGGGTCTGgGgGCTGgGgGATGCTCTGACCTTGAAAGGGTCCCTATCTGGAGAGCG	✓	phdtk-1		
61	CATCCTGCCATTGTGGTTACTGAGTCTGCTGGTTGGACAGAGGAATGGGCAAGCAATGTTCC	✓	pRNdtk-1		
73	AGCGGAGATGCA6CGGCTTGGGGCCACCCCTGCTGTGCTGCTGGCGGCGGCGGCTGCC	✓	phdtk-3		
4	CGAGCGGCA6CGGCGCTGAGAGCGCGCGGCTGCTGGCGGCGGAGAGGAGCGCGCGCGCG	✓	pcdtk-3		
1	✓	pmdtk-2		
1	✓	phdtk-2		
94	GCTCTCTGCA6CCTCCCTCTGCTAGGAGCCAGTGGCCACCTTGAACTCA6TTC	✓	pmdtk-1		
83	AGGAGGTACAAACGTGCTGAAATGTGTGCGGTTCAAGGAGGCAATTTGGTTAAGCCCT	✓	phdtk-1		
121	C6GTGCCCTCTTATTGTCTTTTGGGGYTTTATCTTGGATGGGGGCACTTGGCTTTGTGATGA	✓	pRNdtk-1		

Fig. 2

[illegible]

Fig. 2 (Forts.)

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312	CTGCTA	AAAGCA	ATCATCA	GAAAGT	GAAACC	TGGCAAA	CTTACCT	CCAGCT	ATACACAA	TGAGA	phdtk-3
244	G6GCA	AA	AA	AA	AA	AA	AA	AA	AA	AA	pcdtk-3
106	GCCAC	AG	TC	CC							pmcdk-2
67	GCCAC	AG	TC	CC							phdtk-2
329	GCTCC	AG	TC	CC	AA	6CCG	CG	CA	6CCG	TC	pmcdk-1
314	GCGCT	AG	TC	CC	AA	CCCC	CG	CA	6CCG	TC	phdtk-1
361	GTCAC	AG	TC	CC	AA						prRdtk-1
372	CCAA	CAC	AG	AC	AG	AA	GG	TT	GG	AA	phdtk-3
304	CCAA	CAC	AG	AA	AC	CA	AA	TT	GG	TA	pcdtk-3
148	G6A	AG	AA	AA	AA	AA	AA	AA	AA	AA	pmcdk-2
109	G6A	AG	AA	AA	AA	AA	AA	AA	AA	AA	phdtk-2
389	GA	AA	GC	CA	GG	AA	AG	CG	CG	CG	pmcdk-1
371	G6A	AG	CG	CG	AA	AA	CG	CG	CG	CG	phdtk-1
406	G6A	AG	CG	CA	GA	AA	AG	CG	CG	CG	prRdtk-1
432	T										phdtk-3
364	T	ACA	G	AT	AA	CA	GA	AA	CT	GG	pcdtk-3
208	AT	GG	AA	AT	CT	GG	CA	TC	CC	CA	pmcdk-2
169	AT	GG	CA	AT	CT	GG	TA	TC	CC	CA	phdtk-2
449	AT	GG	AA	AT	AT	GG	CA	TC	CC	CA	pmcdk-1
431	AT	GG	AA	AT	AT	GG	CA	TC	CC	CA	phdtk-1
466	AC	GG	AA	AT	AT	GG	CA	TC	CC	CA	prRdtk-1

Fig. 2 (Forts.)

433
 424
 265
 226
 500
 479
 526
 phdtk-3
 pcdtk-3
 pmcdtk-2
 phdtk-2
 pmcdtk-1
 phdtk-1
 p(RN)tk-1

433
 484
 325
 286
 560
 533
 583
 phdtk-3
 pcdtk-3
 pmcdtk-2
 phdtk-2
 pmcdtk-1
 phdtk-1
 p(RN)tk-1

433
 544
 385
 346
 620
 593
 643
 phdtk-3
 pcdtk-3
 pmcdtk-2
 phdtk-2
 pmcdtk-1
 phdtk-1
 p(RN)tk-1

Fig. 2 (Forts.)

433	phdtkk-3
783	CC TGATGGAGTACTAGAGCGCTGCCCATGTGCAAGTGGCTTGAATCTGGCCAACTCTCAAGAGC	pcdtkk-3
619	AAACACTG.GAAGAGTCAATCAGCAGACTGTGAATTTGTGATTTAAAGCAATTAATGAGC	pmdtkk-2
580	CACCAATTGAAGAACATCAATCAATGGCAGACTGTGAAGTTGTGATTTAAAGCAATTAATGAGC	phdtkk-2
856	...ACCAGACAGTC..TAAATATGATGGACTCTTTTATCTAAATAATGCTACGAAATATC	pmdtkk-1
829	phdtkk-1
883	CGAGGCTACAGAG..CCTGAAAGGACCTTCTCTAAATTAAGCTAAATTAAGACTTTGGTATC	pfRNdtkk-1
433	phdtkk-3
843	AGCCACAGTACTACATCTGTGTGTAACCTGTCCCTCCAAATGAAACCAAGCAAGCAAAACGAA	pcdtkk-3
678	ATGATGGAAACCTGGATTGGAAATGCGGAAGAAATGAGGGATGTGGTAAGAAATGTGGAGGCAAG	pmdtkk-2
640	ATGGTGGAAAAATAAAGGTTCAAGATGCAAGATGCAAGAAAGAAAT...GGCTAAAAATAAAGAAACGTTGAATAAAG	phdtkk-2
910	CTTTATGATTTGTGAGCTCAATCCCAAGGATGTAGGAAATCTTCAAGTGTGTAATTAAGCAT	pmdtkk-1
829	phdtkk-1
941	CTGCAATGTTATTTTCTCAGTTTACATGAAAGTGTCTGTGCTCTTCCCTGAAACCCGGAATGCTG	pfRNdtkk-1
433	phdtkk-3
903	GAAAGATCCCTTGAAACATGGATGAGATGCCCATTTATCAGTTTAAATACCCAGAGATAATCTT	pcdtkk-3
738	AAGAGGCGAGGACTGAAATCAAGTAGAGTCCGACAAACCAACCAAGATCTACCAATGCTTCCG	pmdtkk-2
697	AATATAGATGATCACAAATAAATAAATAAAGATGCCGCGCCGCAAGCTTATTCCTTTA	phdtkk-2
970	TCCGACAAATACTTTCCAAAGCTCTGGAAGTGTAAAGGACTTTGTCTTGTGATGGAACTCCC	pmdtkk-1
829	phdtkk-1
1001	C6CAACTTGTTTCTTTTTTGTGAAGGAACTTCCCTAAATTAATGCTAAATACAGTAAATTAATGCTG	pfRNdtkk-1

Fig. 2 (Forts.)

[illegible]

Fig. 2 (Forts.)

Fig. 2 (Forts.)

433	phdtk-3
1323	CTTTGTACAGCAGAAATAAACGTATCAGTACTCGTACTCAATAAAAAACACACGAGCA	pedtk-3
882	pmdtk-2
769	phdtk-2
1227	pmdtk-1
829	phdtk-1
1298	pRNdtk-1
433	phdtk-3
1383	T	pedtk-3
882	pmdtk-2
769	phdtk-2
1227	pmdtk-1
829	phdtk-1
1298	pRNdtk-1

Fig. 2 (Forts.)

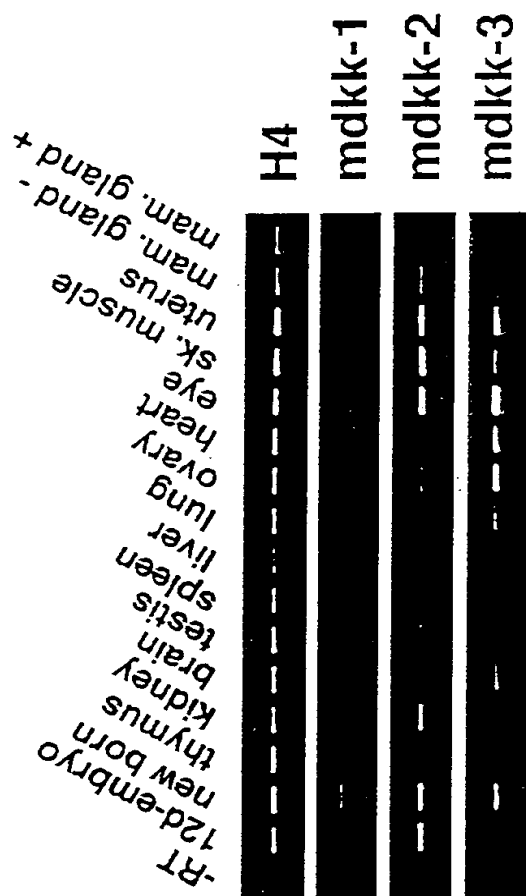


Fig. 3